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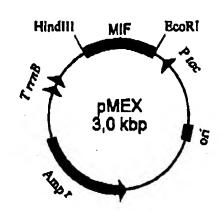
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(54) THE: RECOMBINANT MACROPHAGE MIGRATION INHIBITORY FACTOR: EXPRESSION IN ESCHERICHIA COLI AND PURIFICATION OF RECOMBINANT PROTEIN

(57) Abstract

According to the known cDNA sequence we used the PCR (polymerase chain reaction) technique to isolate a gene encoding human MIF from uterus endometrium The PCR product was cloned in the Escherichia coli plasmid voctor pUC 19 and the nucleotide sequence was confirmed. We were able to successfully express human MIP in the bacteria Escherichia coli by uso of the pKP 1500 expression plasmid. The recombinant protein accumulated intracellularly in soluble form. comprising more than 30 % of total call protein. We have designed an original two step procedure where protein purification was accomplished by gol filtration and ion exchange chromatography. The purified recombinant MIF was used to immunise a rabbit and antibodies obtained were used for MIF detection in human ussues by immunohistochemical techniques.



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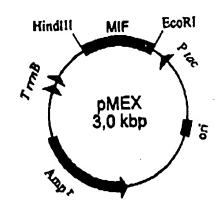
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INTERNATIONAL SEARCH REPORT

Information on passas lamely members

PUI/SI 95/99022

Patent document cited in search report	Publication date	Patent memi	Publication date	
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WO-A-9426307	24-11-94	AU-8- CA-A-	6834594 2163211	12-12-94 24-11-94

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RECOMBINANT MACROPHAGE MIGRATION INEIBITORY FACTOR; EXPRESSION IN ESCHERICHIA COLI AND PURIFICATION OF RECOMBINANT PROTEIN

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BACKGROUND OF THE INVENTION

The macrophage migration inhibitory factor (MIF) was the first lymphokine to be discovered. In 1966 it was shown that antigen stimulated lymphocytes produced a soluble factor that inhibited the migration of macrophages in vitro (Bloom, E. R. & Bennett, B. (1966) Science 153, 80-82 / David, J. R. (1966) Proc. Natl. Acad. Sci. USA 65, 72-77; MIF containing supermatants of stimulated lymphocytes were subsequently shown to alter macrophage functions and enhance the killing of micro-organisms and tumour cells (Churchill, W. E. , Plessens W. F. , Sulis, C. A. , David, J.R. (1975) J. Immunol. 115, 781 / Nathan, C. F., Karnovsky, M.L., David J. R. (1971) J. Exp. Med. 133, 1356 / Nathan, C. F., Remold, E. G., David, J. R. (1973) J. Exp. Med. 137, 275). MIF has also been shown to correlate with delayed type hypersensitivity and deliblar immunity (Bloom, B. R. & Bennett, B. (1966) Science 153, 80-82 / David, J. R. (1966) Proc. Natl. Acad. Sci. USA 65, 72-77 / David, J. R. & David, R. A. (1972) Prog. Allergy 16, 300-449) MIF activity has been detected in the synovia of patients with rneumatoid polyartritis (Odink, K. , Cerletti, N., Bruggen, J., Clerc, R. G., Tarcsay, L., Zwadlo, G., Gernards, G., Schlegel, R. & Sorg, C. (1987) Nature (London) 330, 80-82), in leukocyte culture supernatants of mice during allograft rejection (Al-Askari, S., David, J. R., Lawrence, H. S. & Thomas, L. (1965) Nature (London) 205, 916-917 / Harrington, J. T. (1977) Cell. Immuncl. 30, 261-271; and in various chronic inflammatory loci (Burmeister, G., Zwadlo, G., Michels, E., Brocker, E. & Sorg, C. (1984) Lymphoxine Res. 3, 236 (abstr.), Schlegel-Gomes, R. ,

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Diepgen, T. L., Neumann C., Sorg, C. (1990: Arch. Dermatol. Res. 282, 374-378

However, a purified or aloned lymphckine was required to demonstrate that these altered macrophage functions were induced by MIF and not by any other factor. Low levels of activity empressed by natural sources proved the biochemical characterisation of MIF very difficult, since the native protein has not been isolated in sufficient amounts yet.

Tre uDNA encoding human MIF was cloned in COS cells in 1969 by Weiser et al.. By functional empression cloning of the cDNA from T cells a clone was identified which expressed a strong MIF activity. Waiser also stressed that MIF is the product of activated lymphocytes T only. (Weiser, W. Y., Temple P. A., Witek-Giannotti J. S., Renold, H. G., Clark, S.C. & David, J.R. (1989) Proc. Natl. Acad. Sci. USA 86, 7522-7526). Supernatants from COS cells :bearing recombinant gene encoding human MIF) were shown to stimulate the antibody synthesis (Weiser, W. Y., Pozzi, L. M., David U. R. & Titus, R. G. (1992). Proc. Natl. Acad. Sci. 39, 6049studies Preliminary Ofi Leischmania intracellular parasite; infections (Weiser, Y. W. , Ponzi, 1. M. & David J. R. (1991) J. Immunol. 147, 2056-2011) and the appointmention that MIF containing supernatants from recombinant COS cells have the ability to activate macrophages to express mitric oxide synthetase and to produce NO (Cunta, F. Q., Weiser W. Y., David, J. R., Moss D. W., Moncada, S. & liew, F.Y. (1993) J. Immunol. 150, 1908 -1912 / Liew, F. Y. Miliot, S., Farkinson, C., Faimer, R. M. J., Moncada, S. (1990) J Immunol. 129, 351) indicated that the newly discovered (12 kDa) protein probably acts as a critical factor in cell mediated immune host defences . Recently it was reported that MIF plays a central role in the toxic response to endctomemia and possibly toxic shock (Bernhagen, J., Calandra, T., Mitchell, R.A., Martin, S. B., Tracey K., J., Voelter W., Manogue K. R., Cerami, A. & Bucala R. (1993) Nature 365, 756-759: MIF was isolated from ocular lens by Wistow et al. in 1993 (Wistow G. J., Shaughnessy, M. F., Lee,

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D. C., Hodin, J. & Zelenka, P. S. (1993) Proc. Natl. Acad. Sci. USA 90, 80490-80529) and a putative MIF from rat liver was isolated by Blocki et al. in 1992 (Blocki, F. A., Schlievert, P. M. & Wackett, L.F. (1993) Nature 360, 269-270). This protein that matches the primary structure of a human leukocyte MIF (Weir, 1989) in 25 out of 26 terminal amino acids and which has both glutations transpherase and MIF activity was shown to link chemical and immunological detoxilication systems. Suzuki and colleagues (Suzuki, M. , Murata E. & Tanaka I. (1994). C. Mol. Biol. 235, 1141-1143; crystallisation preliminary and tuods reported crystalle traphic studies of MIF from human tymphocytes. From this art. le it can be realised that human MIF was expressed in E. coil and purified by affinity chromatography on the basis of the results of Block! et al. (Block!, F. A., Schlievert, P. M. & Wackett, L.P. (1992). Nature 360, 269-270;

Although the focus of research and hence the need for this process continues to expand and despite the fact that MIF from natural sources has never been isolated in sufficient amounts yet and that the quantity of recombinant protein. produced by tissue cultures is low, the information about empression of MIF in bacteria are very scarce. The data about the yields are generally not reported and the purification procedures are only mentioned and not well described. Some purification procedures include affinity chromatography based on glutathione transferase activity of MIF. The identity of purified proteins, isolated by affinity chromatography, was however not checked by N-terminal aminc acid sequencing although it should be since glutathione transferase activity for human MIF and confirmed controversial. In addition, the above mentioned affinity chromatography procedure could not specific be recombinant protein (bacterial proteins with glutathione affinity could also be isolated).

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On the other hand, as we know, at MIF protein is commercially available, although human cDNA is marketed by some companies.

SUMMARY OF THE INVENTION

The present invention relates to the construction of an expression vector for human MIF in Escherichia coli and the establishment of an original optimised protocol that enable to express and isolate large amounts of highly purified and biologically active recombinant numan MIF. From 50 grams of recombinant bacterial cells (wet weight) 1 gram of highly purified and biologically active protein can be obtained. In this way various biochemical, biophysical and physiological studies of this still poorly understood cytokine should be facilitated. MIF could have also specific therapeutic and diagnostic values. From this point of view it is still more important to obtain it in large quantities and in a pure, biologically active form.

Purified recombinant MIF could serve to produce MIF specific antibodies which could be used to detect MIF in various human tissues by immonistochemical methods. We clearly demonstrated the presence of MIF in epithelial cells of some samples of human uterus endometrium (where it is probably connected with certain pathologic states), in epithelial cells of various inflammatory tissues (human and animal) and in the same cell type in allograft rejections of human kidney. Kidney epithelial cells were almost completely negative in healthy individuals.

In this way MIF and (or! against MIF directed antibodies could be used as diagnostic or prognostic markers.

The present invention thus relates also to MIF that is expressed in epithelial cells. Besides its general function in the immunological system, the above described discoveries opened also the possibility that MIF acts as an important factor in mucosal immunity. Epithelial cell MIF could be a

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signal for injury, infection, tissue invasion, ect. and could serve as a chemoatrantant and activator of macrophages (and possibly other immune cells) on the site of inflammation. In this sense it could serve as one of the first cytokines in the complex cytokine network.

MIF alone or in combination with other moleculas (such as IFN-Y, IL-2, ect.) could be used to treat diseases where cellular / mucosal immunity should be stimulated (such as infections, AIDS, cancer, etc.). On the other hand, Frainst infections antibodies or MIF antagonists could be use; when immunological functions should be scaled down (like in the immunological functions should be scaled down to organizes of auto immune diseases, tissue or organizations, etc.).

The present invention is thus directed to purified recombinant MIF, to its analogues, MIF specific antibodies and antagonists that could be (in any pharmaceutical combination) used as diagnostic, prognostic or therapeutic markers/agents.

DETAILED DESCRIPTION OF THE INVENTION

Amplification by polymerase chain reaction (PCR) on a cDNA template could be used to isolate the MIF coding region and to create restriction sites which enable to express MIF in vectors. coli expression Escherichia different ge! analysed could be products amplification electrophoresis. Vectors, such as pKP 1500 could be used to express the recombinant protein in the cytoplasm of E. coli cells. Cthers, such as pIN-III-c p A2 (containing a signal sequence) could be used for transporting MIF into the E. col: periplasmic space.

periplasmic space.

The efficiency of the expression due to different expression vectors (and E. coli strains) could be determined by protein analyses of whole cell extracts (lysates). The cell extracts (lysates) are prepared as described in example 1 and analysed by SDS-PAGE and LEF using known mol. wt. or pl

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standards and a control to total cell typate of crude extract of nost E. coll cells, bearing the expression vector without insert).

A strong new band (corresponding to an approximately 12 kDa protein) appears at the expected position when 5, cold cultures with recombinant pKP 1500 plasmid, harbouring MTF insert are analysed.

According to the results of SUS-PAGE E. coll YM 109 pearing the recombinant pKP 1500 plasmid (designed pMEM) and empressing MTF intracellularly could be selected for large scale production (fermentation) and subsequent purification of the protein. All purification steps should be done at 400 to minimise proteolysis. Sel filtration chromatography could be used as a first purification step. The aim of this choice is to remove very large molecules (nucleic acids such as plasmids and chromosomal DNA, high molecular proteins, etc.

and very small ones (such as many bacterial toxins and pyrodens, culture media residual ingredients, ect.). On the other hand eventual scluble multimeric molecules, resulted of improperly folded recombinant protein, could be eliminated in this way. Purification of the product could be monitored by SDS-FAGE and IEF that produced bands at expected positions according to the known mol. wt. and pl standards and in the comparison with the control. Fractions which are found to be reach in recombinant protein are pooled, concentrated and further purified by ion exchange chromatography. Different buffers at different pH could be used at this purification step. A successful approach is when 10 mM phosphate buffer (pH 6,4) is used in combination with a CM-cellulose ion exchange column. This has also a logical explanation since the isoelectric point (pI) of recombinant human MIF is approximately 7,0 and bacterial proteins are predominantly more acidic. By using the phosphate buffer mentioned above it could happen that the recombinant protein did not bind to the CM cellulose. In this case it is recommended to adjust the pH

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very carefully college lower the pF for 0.1 or 0.2 units. The elution diagram of the purification by classical cation exchange chromatography on a CM cellulose column, using phosphate buffer pR 6.4, shows two peaks only. The first one represents unbound impurities (without recombinant protein), the second one is highly purified recombinant numan MIF.

We have successfully expressed the human matrophage migration inhibitory factor (MIF) in Esamerichia coli by use of the pME 1500 expression plasmid containing the tac promoter and a temperature sensitive origin of replication, ensuring a high plasmid copy number at elevated temperatures. The recombinant protein accumulated intracellularly in soluble form, comprising more than 30% of total cell protein. have designed a two step procedure where protein purification was accomplished by gel filtration on Sephaces G-50 and cation exchange chromatography on CM cellulose columns. The 12 kDa protein was shown to be pure by SDS-PAGE, IEF and by HPLC. The identity of the purified protein was verified by amino acid analyses and N-terminal sequencing. The MIF assay was used to measure its activity. I gram of highly purified and biologically active recombinant human MIF was obtained from 50 grams of E.coli cells (wet weight.. CD spectra in the near UV and NMR analysis confirmed an ordered, native like structure of purified recombinant MIF.

During developing the purification procedure attention was given particularly to two problems. Firstly, isolation of soluble recombinant proteins from whole cell lysates as opposed to isolation of inclusion bodies or proteins from periplasmic space is more laborious. It is difficult to remove bacterial debris by filtration or centrifugation que to high viscosity. In this case the disruption of cells by French Press was proved unsuccessful. If cells are only partially broken by freezing and thawing only, following the procedure described in example 1, problems with viscosity are overcomed and this represents also a first purification step.

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The socora problem are the activity measurements. MIF assay is a biological test and hence laborious, time consuming and with lower reproducibility. On the other hand many factors (such as impurities in the sample) could contribute to the results of the test. So the activity measurements are not a good shoice for MIT detection during the purification procedure. The pl of the recombinant MIF (approximately 7,0) is very different in comparison to bacterial proteins which are predominantly acidic (with pH values below 6,4). After isoelectric focusing of a control (total protein extract of E. poli YM 109 pells bearing the empression vector pRF1500 without insert; there are no bands at the expected position (pl higher than 6,4). This fact and excellent expression of the recomplinant MIF permit the design of a purification procedure where all steps could be followed on the basis of IEF analyses of fractions without activity measurements.

We would like to stress that the excellent expression is not due only by the expression vector chosen (pKP 1500), but it is also the result of bacterial ceil cultivation. Due to cultivation conditions the cells could be kept as long as needed in a highly repressive state regarding the production of recombinant MIF. The way of the preparation of the seed culture could be in this case very important since when growing in certain complex media, the cells could produce the recombinant protein even if they are not induced. Derepression during cell growth can have negative influences on biomass production and hence on the final yields of recombinant protein.

The procedure is specially adopted for large scale production and purification using classical methods. However it does not mean that quick purification methods with small capacities (such as FPLC) could not be used. If the concentration of endotoxins in the end product (purified MIF) is too high, additional purification steeps (such as reverse-phase chromatography) could be used. However, this is not necessary if the roles of aseptic work are respected.

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Purified recombinant MIF served to produce MIF specific antibodies which we used to detect MIF in various human tissues by immunchistochemical methods. We clearly demonstrated its presence in epithelial cells of some samples of human uterus endometrium (where it is probably connected with certain pathologic states), in epithelial cells of various inflammatory tissues (human and arimal) and in the same cell type in allograft rejections of numan kidney, kidney epithelial cells were almost completely negative in nealthy individuals.

EXAMPLE 1

Construction of expression vectors

The coding region of MIF DNA was isolated by polymerase chair, reaction (PCR). Double stranded cDNA that was used as a template for PCR was synthesised using mRNA from human uterus endometrium or from epithelial origin cultured cells and chemicals supplied by Amersham (cDNA synthesis kit). The reactions were done according to the Amersham instructions, mRNA was isolated by affinity chromatography on oliqo dt spun columns (Sandrook, J., Maniatis, T. & Fritsch, E. F. Molecular Cloning: A laboratory manual; Cold Spring Harbor Laboratory (1989). The guanidinium izothiocianate method followed by isopicnic ultra centrifugation in a gradient of caesium trifluoroacetate was used for total RNA isolation.

The primers used were:

I.: 5' GGATCCGAATTCATGCCGATGTTCATCGTAAACACCA 3' (sense strand oligonucleotide corresponding to the N terminus of MIF; Eco RI cutting site underlined and start codon in bold)

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II.: 8' GTCGACAAGCTTTTAGGCGAAGGTGGAGTTGTICCA 3'
(antisense strand oligonucleotide corresponding to the C terminus of MIF; Hind III cutting site underlined and stop codon in bold)

The same primers in the same PCR were used to create. Eco RI and Hind III cutting sites. Flanking the coding region that allowed directional cloning into multiple cloning sites of D. coli vectors pUC 19 (Messing, J. (1983) Methods in Enzymology, Vol. 101, pp. 20-78 (Wu, R., Grossman, I. 6 Moldave, K., Ed.). Sar. Diego:Academic Press), pIN-III-ompAl (Auerswald, E. A., Genenger, G., Mentele, R., Lenzen, S., Assiery-Machieidt, I. Mitschang, L., Oscherner, H. & Fritz H. (1991). Eur. J. Biochem., 200, 131-156 / Ghrayeb, J. , Kimura, H. , Takahara, M. , Hsiung, H., Masui, Z. , Inouye, M. (1964) EMBO J., 3 (10), 2437-2442) and pKP 1500 (Miki, T., Yasukochi, T., Nagatani, H., Furuno, M., Orita, T., H., Imoto, T., Horiuchi, T. (1987) Protein Yamada, 327-332). Oligonucleotide synthesis was Emgineering 1, carried out using an Applied Biosystem DNA synthesiser, according to the manufacturer recommendations. The products were cleaved from columns and de protected by saturated ammonium hydroxide and further purified by polyacrylamide gel electrophoresis.

The cycling conditions for the polymerase chain reaction were denaturation at 94°C for 1 minute, annealing at 55°C for 1 minute and extension at 72°C for 2 minutes. 100 μl of the reaction mixture contained: 10 mM Tris (pH 9.4), 50 mM KCl, 100 ng cDNA (incubated in a boiling water bath for 10 the reaction mixture),1 added to minutes prior oligonucleotide I, 1μ M oligonucleotide II, deoxynucleotide triphosphates (dATF, oCTP, dGTP dTTP; 200 μ M concentration each), ImM MgCl2, 2,5 units of Tag polymerase (Perkin Elmer) : 100 µl of mineral oil was added on the top to prevent evaporation. A total of 30 PCR cycles were carried out in an Perkin Elmer thermo cycler. Terminal extension was allowed by heating the PCR product at 72°C for 10 minutes. Mineral oil was then carefully removed and following fenolisation and

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echanol precipitation (Sanbrook, C., Maniatis, T. & Fritsch, E. F. Molecular Cloning: A laboratory manual; Cold Spring Harbor Laboratory (1989), restriction of the approximately 400 base pair fragment with EccRI and Hind III endonucleases was performed for at least five hours under standard conditions (Sambrook, J., Maniatis, T. & Fritsch, E. F. Molecular Cloning: A laboratory manual; Cold Spring Harber Laboratory (1989). The reaction mixture was then again fenolised. DNA was ethanol precipitated, dried in a speed-vac mondentrator, resuspended in 50 µl of TE buffer and purified by get filtration chromatography (1 ml Sephacryl S-300 spin columns. The E. coli plasmid vectors were digested and purified in the same way; except that Sephacryl S-400 was used in the gel filtration chromatography step. Inserts were ligated to vectors in a molar ratio 3 :1 in favour of inserts and the ligation reaction was performed at 1500 for 15 hours under otherwise standard conditions

A fluorescent method for quantifying mg amounts of DNA was used to measure the concentration of vector and insert DNA (Sanbrook, J., Maniatis, T. & Fritsch, E. 7. Molecular Cloning: A laboratory manual: Cold Spring Harbor Laboratory (1999)). I µl of the ligation mixture was used to transform E. coli DH5α competent cells (Earnahan, D. (1983) Mol. Biol. Techniques (1985) Hannahan, D. 557-580, E.coli. In: DNA cloning: a practical uransformation of arroach, Vol.1, pp109-135 (Glover, D. M. Ed.). Oxford: IRL Press;. Eighteen of the resulting clones were picked up and amplified in 3 ml of LBA medium. Plasmids were then isolated (Del Sal, G.; Manfioletti G. & Schneider C. (1989) Nucleic Apid. Res. 16, 9878) and MIF positive clones were identified restrict home analyses. The nucleotide sequences of four positive clones from three independent PCR were confirmed by the dideoxy sequencing method (Sanger, F., Nicklen, S. & Coulson, A. R. (1977) Froc. Natl. Acad. Sci. USA. 74, 5463]. The nucleotide sequence and the deduced amino acid sequence of the sequenced clones are shown below

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ATG	CCG	ATG	170	VIC	ATC	ARC	ACC	AAC	GTG	ccc	ce:	ودد	тсс	GTG	45
Ke:	Pre	Met	Phe	1.0	Và l	Asn	Thr	Asn	Val	Pra	Arç	Ala	Ser	Val	
				•					10					15	
ccs	SYC	G55	הזה	crc	TCC	GA:3	ctc	ACC	CAG	CAG	CTG	SEC	CAG	SCC	90
ero	Asp	Gly	Phe	Len	Ser	۳. C1	Leu	Thr	G_ n	GIr.	Let	γĮο	Gln	λia	
				20					25					30	
A22	561	۵۸۵	ccc	ccc	CAG	TAC	λτς	ccc	GT G	CYC	STS	G1:	ccs	GAC	135
Thr	617	Lys	Fro	Fr:	Sin	Ty:	lie	817	va:	H13	٧ė.	127	f:o	AFD	
				35					÷0					45	
CAS	crc	ATG	GCC	TTC	GGC	GGC	TCC	AGC	GAG	ccs	TSC	600	ctc	7G::	190
G' n	Leu	Met	A:a	?he	Gly	Gly	Ser	Ser	G1	Pro	Cys	Ala	Leu	Cyz	
				50					55					60	
AGC	276	CAC	AGC	ATC	GCC	AAG	ATC	GGC	GGC	GCS	CAG	AAC	CGC	TCC	225
Ser	Leu	Hie	3er	Ile	G) y	Lys	11e	G) y	Gly	hla	Sin	<u>As n</u>	Arg	Ser	
				65					0 5					75	
TAC	λGC	AAG	CTG	CTG	tec	GGC	CTG	CTG	GCC	GAG	CSC	CTG	೦೯೭	ATC	270
7yr	Sec	Lys	Leu	Leu	Cys	GIY	Leu	Leu	Aia	Glu	3:0	Leu	۸:۵	11e	
				80					85					90	
								•							
AGC	ccs	GAC	AGG	GTC	TAC	ATC	AAC	TAT	TAC	GAC	ATG	AAC	GCG	GCC	315
Ser	Pro	Arp	Arg	val	Ty:	lle	Asn	Tyr	Tyr	Asp	Met	Aen	Ala	Ala	
				95					100					105	
AAT	GTG	GGC	TGG	AAC	AAC	TCC	ACC	TTC	GCC	TAK					360
Asn	Val	Giy	Trp	Asn	λzn	Ser	Thr	Phe	Als	•					
		•	•	110					:15						

With the exemption of one codone only, resulting in all cases in the same amino acid change, the sequences were identical as reported (Weisser, W. Y., Temple F. A., Witek-Giannotti J. S., Remold H. G., Clark S.C. & David J.R. (1999)

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Proc. Natl. Acad. Sci. USA 86, 7522-7526; The amine soid substitution was exactly the same as found by Wistow (Wistow G. J., Shaughnessy, M. F., Lee, D. C., Hodin, J. & Zelenka, P. S. (1993) Proc. Natl. Acad. Sci. USA 90, 8049-30529). At. this moment in the literature exists a great confusion. between cDNA/proteins with very similar nucleotide/amino acid sequences and between the biological functions these aDNA products/protoins could express.

One of the recombinant puC 19 plasmids, harbouring the MIP insert was Eco RI and Hind III digested and fractionated by get electrophoresis. The band containing DNA of 400 page pairs was excised and isolated from the gel slice by adherence to glass powder (Vogelstein, S. & Gillespie, D. (1979) Proc. Natl. Acad. Sciu. USA 76, 615-619). The MIF coding region was then subcloned into empression vectors pIN-III-ompA2 and pKP 1500 following the same procedure as described above.

Empression and purification

Step 1: Fermentation conditions

200 ml of saturated culture of E. coli YM 109 strain (Miller, H. (1987) Methods in Enzymology 152,145-170), harbouring the MIF expression plasmid pMEX (pKP 1500 with MIF insert was used to inoculate a 15 l (Chemap LF 7/14/20) bicreactor containing 10 l of sterile LB broth (10g tryptone, 5g yeast extract and 10g of NaCl per 1 of media) with 10 ml of Silicon 1510 (added as anti foaming agent prior to autoclaving). The seed culture was prepared by inoculating 10 μi of frozen glycercl stock culture into 200 ml of M9 minimal liquid medium, supplemented with 100 mg ampicillin per litter (Sanbrook, J., Maniatis, T. & Fritsch, E. F. Molecular Cloning: A laboratory manual; Cold Spring Harbour Laboratory (1989)) in an 500 ml Erlenmeyer flask and by cultivating at 21°C for 36 hours with agitation on a rotatory shaker {140 rpm:. Following seeding, ampicillin (100 mg per 1) was added

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to the growth medium asseptically. Formentation was carried out with controlled stirring (600 rpm) and aeration (301/min). The culture was incubated at 37^{9}C for 2,5 to 3 hours (until it reached an A600 of 0,5 to 0,1) after which it was induced with IPTG (0.59/10 l culture) and cultivated for another four hours under the same conditions.

Step II: Preparation of cell entract

Four hours after induction with IPTG, the bioreactor was water cooled to 120C, the culture decented into glass flasks immediately chilled on ice. Bacteria from fermentation broth were harvested by centrifugation at 8000g and 40% for 20 minutes. A total of 50 grams of cell pellet was suspended in Zoo mi of sterile water. The suspension was freeze-thawed three times and then sonicated with one burst (1,5 minute) in a 70 W ultrasonic ice water bath. Following centrifugation at ECOO g for 20 minutes at 400 to remove cellular debris the supernatant was filtered trough a Millipore low protein binding membrane (C, 2 un) transferred to a 350 ml stirred ultrafiltration cell (Amicon) equipped with YM2 (2000 mol. wt. cut off) membrane and concentrated five fold at 4°C.

Step III: Gal filtration chromatography

A glass column (50 mm diameter) was packed with Sephadex G-50 (bed height 1500 mm) and equilibrated with buffer A (0,1 M Tris buffer pH 7,4; 0,3M NaCl; 1 mM EDTA). The sample (20 ml of the concentrated supernatant from step II) was carefully added on the top of the gel bed and eluted with buffer A mentioned above. Fractions were collected automatically. The flow rate was 42 ml/h and the fraction volume was 14 ml. The total procedure was performed at 4°C and it required approximately 32 hours. The Sephadex G-50 fractions were monitored by absorbance at 280 nm for protein. Some fractions were analysed by SDS-PAGE and IEF where the protein extract of E. coli YM 109 cells, bearing the expression vector without insert, was used as a control. Aliquots of fractions were dialysed by a mini dialysis method (Francky Andre); personal communication) prior to further analyses by IEF and

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SDS-PAGE. Several fractions containing recombinant protein were pooled and concentrated on Amicon concentrator to 150 ml.

Step IV: ion exchange chromatography CM ceilulose which had been previously regenerated according to the manufacturer instructions and suspended in buffer B (10 mM phosphate buffor/pH 6,4) was used to pack the column which was then equilibrated with the same buffer. 75 ml of the concentrated sample (from step III) was dialysed against buffer B and loaded on the CM cellulose column. (bed volume 250 ml.. After the column was washed to remove unbound protein (until the absorbance at 280 nm was less than 0,05) the protein was eluted with a linear NaCl gradient generated from 500 ml of buffer B and 500 ml of buffer E containing 0,3 M NaCl. The separation was performed at $4^{\rm OC}$. The eluate was collected automatically with a flow rate of 19 ml/m. Fractions of 6,3 ml were collected. Elution was monitored in the same way as described above. All fractions from the second peak contained purified recombinant MIF. They were dialysed against distilled water at 400 concentrated to a concentration of 5 mg per ml ultrafiltration. The purified protein was then stored at minus 70°C . Some samples were lyophilised prior to storage.

Electrophoresis

Protein purity and molecular weight were determined by abdium dodecyl sulphate polyacrylamide electrophoresis (SDS-PAGE) on a Pharmacia Phast System using prefabricated 8 to 25 % gradient polyacrylamide gels and the Sigma SDS 7 molecular weight standard mixture (containing seven proteins in the 14200 to 94000 mol. wt. range). The proteins were electrophoresed and stained with Comassie brilliant blue G-250 according to the manufacturer instructions.

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For protein analyses of whole cell lysates, cells were pereted by centrifugation, resuspended in water and mixed with an equal amount of 2% SDS loading buffer, boiled for 5 minutes and sonicated with one burst (1 minute) in an ultrasonic water bath. After centrifugation (Eppendorf microcentrifuge, 14000 rpm, 10 minutes) the supernatant was diluted with 1% loading buffer when necessary and subjected to SDS -FAGE.

Isoelectric facusing (IEF)

An IEF gel (thickness 1 mm) was prepared using 5% T and 5% C, 10% glycerol, 0.45 % ammonium persulphate and 6.66% Tharmalytes (pH 3-10). Prefocusing was carried out at a constant power of 25 W (voltage limit 300 V) at 7°C. After the samples (20-50 µl each), together with standards (pl 3.5 to 9.3 / Pharmacia), were spotted on the gels, the IEE was carried out at constant power of 25 W (voltage limit 1500 V) for approximately 2 hours. The proteins in the gel were lixed by soaking the gel in 20% trichloroacetic acid for 30 min; stained with a solution of 0.2% Coomassie blue G-250 / 45% methanol / 10% acetic acid for 5 min and destained with the same solution without G-250. Alternatively, proteins were focused on a Phast Gel System (Pharmacia) using precast gels, pH range 3 to 9.

Determination of the N-terminal aminoacid sequence

The amino terminal sequence of the recombinant protein was determined by Edman degradation using an automated Applied Biosystems (Foster City, CA) Model 477A pulsed liquid phase protein sequencer with an Model 120A on line FTH amino acid analyser. Sequencing was performed with regular cycle programs and chemicals from the manufacturer.

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MI: ocological assay

The biological activity of the recombinant human MIF was determined using guinea pig peritoneal macrophages as indicator cells according to the method described by Harrington et al. (Harrington J. T., JR. & Stastny F. (1973), J. Immunol. 110 (3), 752-759).

The percentage of inhibition was derived as follows: % inhibition = 100 - (average migration of test samples/average migration of control samples) X 100. Inhibition of 20% or greater was considered to be significant

<u>lmunohistochemical methods</u>

For immunohistochemical detections of MIF we used commercial kits and we followed the instructions of the manufacturers. We used primary antibodies developed against purified recombinant numan MIF.

The above described example serves just for illustration and do not limit the scope of the present invention.

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CLAIMS

- 1. Oligonucleoride primers (their sequence is shown below) and oligonucleorides with other sequences, characterised by the ability to enable amplification of the nucleoride sequence, encoding human MIF by polymerase chain reaction and subsequent cloning of the PCR product in different vectors.
- 1. : 5' GENTOCGARTTCATGCCGATGTTCATCGTAAACACCA 3'
- 11.: 8' STCGACAAGCTTTTAGGCGAAGGTGGAGTTGTTCCA 3'
- 2. The expression vector construct pMEX, characterised by including the below presented nucleotide sequence, encoding human MIF or by including other nucleotide sequences, encoding other proteins which are not different in more than twenty amino acids from the sequence described below and that all sequences mentioned are integrated into a vector on the way shown on fig. 1.
 - ATG CCG ATG TTC ATC GTA AAC ACC AAC GTG CCC CGC GCC TCC GTG 45

 Met Pro Met Phe Ile Val Asn Thr Asn Val Fro Arg Ala Ser Val

 S 10 15
 - COG GAC GGG TTC CTC TCC GAG CTC ACC CAG CAG CTG GGG CAG GCC 90

 Pro Asp Gly Pne Leu Ser Glu Leu Thr Gln Gln Leu Ala Gin Ale

 20 25 30
 - ACC GGC AAG CCC CCC CAG TAC ATC GCG GTG CAC GTG GTC CCG GAC 135
 Thr Gly Lys Pro Pro Gin Tyr 11e Ale Val His Val Val Pro Asp
 35 40 45
 - CAG CTC ATG GCC TTC GGC GGC TCC AGG GAG CCG TGC GCG CTC TGC 180

 Cln Leu Net Ala Phe Gly Gly Der Ser Glu Pro Cys Ala Leu Cys

 50 55 60

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ACC CTG CAC AGE ATC GGC AAG ATC GGC GGC GCG CAG AAC CGC TCC Ser Leu His Ser lle Gly Lys Ile Gly Gly Ala Gin Asn Are Ser THE AGE AME CTG CTG TGC GGC CTG CTG GGC GAG CGC CTG CGC ATC Tyr Ser Lys Lau Lau Cys Cly Leu Leu Ala Giu Azo Leu Aro Ile 90 AS. AGO DOG GAD AGG GTO TAC ATC AAD TAT TAC GAD ATG AAC GCG GCD Ser Pro Amp Arg Val Tys lie Amn Tyr Tyr Amp Met Amn Ala Ala 100 360 AAT GTG GGC TGG AAC AAC TCC ACC TTC GCC TAA Amn Vol Gly Trp Amn Amn Ser Thr Phe Ala . 115

The procedure of recombinant E. coli cultivation characterised by such a combination of expression vector, pacterial host strain and cultivation conditions that recombinant MIF is produced in soluble form in the heterologous system of Escherichia coli

:10

- The procedure of isolation and purification of recombinant MIF characterised by the combination of following methods:
- a.) the method for bacterial cell disruption by freezing and thawing only
- b.) two step pusification protocol where gel filtration and ion exchange chromatography are included
- c.) the method for MIF quantification/estimation during the purification procedure on the basis of IEF, SDS-PAGE, Western blotting or ELISA analysis

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5. The below presented nuclectide sequence characterized by encoding human MIP in the epithelial cells of different tissues, like in epithelial cells or the human uterus encometrium.

473	Q0 6	ATG	355	ATC	GTA	AA:	AIC	anc	3T.3	CEE	COL	CCC	100	4:4	4,5
5// J	375	622	710	CTC	rec	GA/J	ere	ATT	ÇAG	UNG	C. G	GCG	CAG	GCC	90
ACC	esc	AAG	cca	SSS	CAG	TAC	λτς	GCG	GTS	CAC	GT.G	670	೧೮೮	SAC	135
CAS	270	ЛTG	655	TTC	GGC	GGC	TCC	AGC	SAS	ccG	TGC	GCG	CTC	TGC	186
NGC	c:6	CAC	AGC	ATC	GGC	arg	ATC	GGC	GGC	GCG	CAG	AAC	CGC	TCC	225
IAI	AGC	AAG	CTG	CIG	TGC	GGC	cts	CTG	ecc	GAG	CGC	CTG	CGC	ATC	270
AGC	cçG	GAC	AGG	GIC	TAC	ATC	AAC	TAT	TAC	СУC	ATG	AAC	ccs	GCC	315
aat	STG	ಽಽಽ	:56	AAC	AAC	700	ACC	710	GCC	aat aat					360

6. The recombinant protein, characterised by the below presented amino acid sequence or other amino acid sequences that do not differ in more than twenty amino acids from the sequence shown below and which is obtained on the way described (related to the points 3 and 4) or on any other similar manner and its therapeutic, diagnostic and any other uses which are not for scientific purposes only.

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lte i	CIO	Ket	Pite	11e 1	Val	A¤≀ı	Thi	Yau	Val 10	Pro	ATĢ	Ala	Sez	Val 15
Pře	A S≨	Gly	Phe	Le ú 20	Ser	61 u	Leu	Thr	61r 25	SIn	Leu	Ala	51:.	A) # 30
Thr	Siy	Lys	?r>	Frc 35	Gl f.	туr	lie	Ala	Va1 40	ein	Va)	Val	Pro	Лер 45
Gl a	Low	Mat	λla	the 50	G1 y	Giy	Ser	Se:	G1u 55	Pro	Суз	Ala	Leu:	Cys 00
Ser	Leu	Ilis	Ser	11e 65	Gly	Lys	lie	Gly	Gly 70	λia	Gln	<u>A=n</u>	λεψ	Se 1
Tyt	Sei	Lys	Leu	Leu 30	Суз	GJ Y	Leu	Leu	A) a 85	G: u	λ'nφ	Leu	Ara	ñJ آاه
>er	٢٢٥) Asp	λrą	V&1 95	1yr	11e	- Авп	туг	Tyr 100		Met	. Asn	414	Ala :05
). en	, Va;	G; A	Csp	<u> </u>		391	The	Phe	115					

7. Antibodies (policional or monoclonal) which are prepared by use of the recombinant MIF (related to the point 6) and their therapeutic, diagnostic and any other uses that are not for scientific purposes only.

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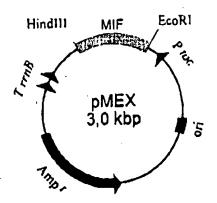


Fig. 1



INTERNATIONAL SEARCH REPORT

Inu onal Application No PCI/SI 95/00022

A CLASSIFICATION OF JUBLECT MOTTER A61K39/395 C07K16/24 C12N1/21 A61K38/19 C12N15/19 C12N15/70

According to International Patent Clarefleation (IPC) or to both national dismilication and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by distrification symbols) IPC 6 CO7K C12N A61K

Documentation provided other than minimum documentation to the extent that such documents are included to the fields statisfied

Electronic data base consulted during the international search (parts of data base and, where practical, jearch terms used)

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Y	claims; table i	
Y	PROTEIN ENGINEERING, vol. 1, no. 4, 1987 ENGLAND GB, pages 327-332. TAKEYOSHI MIKI ET AL 'Construction of a plasmid vector for the regulatable high level expression of eukaryotic genes in Escherichia coli : an application to overproduction of chicken lysozyme' cited in the application see page 328, right column; figure 1	2,3
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X Further documents are listed in the continuation of box C.	Patent family members are listed in onnes.
"Sprow categories of died documents: "A" document defining the general rate of the art which is not considered to be of paracelar relevance. "E" satist document but published on or after the international filing date. "L" document which may throw doubts on priority dainful or which is and to insubitish the publication date of another alabon or other thanks trained (as specified). "O" document referring to an orisi disclosure, time, schubition or other means. "P" document published prior to the international filing date but layer than the priority date claimed.	"I later document published after the externational filité date or procesy date and not in conflict with the application but cited to understand the practice or theory undersying the invertion. "X" document of particular relevance; the clumest invention survoire an inventive stop when the document at the atoms involve an inventive stop when the document is taken atoms carried to considered to involve an inventive stop when the document is committed to invention an invention of particular relevance; the dalened invention carried to considered to involve an inventive stop when the document is combined with one or more other such document, such combined with one or more other such document, in the art. "A" document member of the same patent (arrely)
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Name and mailing address of the ISA Buropean Patent Office, P.B. 5218 Pentitional 2 NL - 2210 HV Running Tel. (* 31-70) 340-2040, Te. 31 611 epo al, Fax: (* 31-70) 140-3016	Le Cornec, N

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	of a cDNA encoding a numan matrophage Nigration Inhibitory Factor' cited in the application	
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	see example / see page 54, line 30 - page 58, line 3	2
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